

MLLNVLRICI IVCLVNDGAG KHSEGRERTK TYSLNSRGYF 40
 RKERGARRSK ILLVNTKGLD EPHIGHGDFG LVAELFDSTR 80
 THTNRKEPDM NKVKLFSTVA HGNKSARRKA YNGSRRNIFS 120
 RRSFDKRNTE VTEKPGAKMF WNNFLVKMNG APQNTSHGSK 160
 AQEIMKEACK TLPFTQNIVH ENCDRMVIQN NLCFGKCISL 200
 HVPNQDRRN TCSHCLPSKF TLNHLTLNCT GSKNVVKVVM 240
 MVEECTCEAH KSNFHQTAQF NMDTSTTLHH 270

Figure 1. Deduced amino acid sequence of *Xenopus cerberus* protein. SEQ ID NO:1.

000001-03101

DO NOT WRITE IN THESE SPACES

GAATTTCCCGAG	CAAGTCGCTC	AGAAACACTG	CAGGGTCTAG	ATATCATACA	ATGTTACTAA	60
CTTAAGGGTC	GTTTCAGCGAG	TCTTTGTGAC	GTCCAGATC	TATAGTATGT	TACAATGATT	
ATGTACTCAG	GATCTGTATT	ATCGTCTGCC	TTGTGAATGA	TGGAGCAGGA	AAACACTCAG	120
TACATGAGTC	CTAGACATAA	TAGCAGACGG	AACACTTACT	ACCTCGTCC	TTTGTGAGTC	
AAGGACGAGA	AAGGACAAAA	ACATATTCAC	TTAACAGCAG	AGGTTACTTC	AGAAAAGAAA	180
TTCTGCTCT	TTCTGTTTT	TGTATAAGTG	AATTGTCGTC	TCCAATGAAG	TCTTTTCTTT	
GAGGAGCAG	TAGGAGCAAG	ATTCTGCTGG	TGAATACTAA	AGGTCTTGAT	GAACCCACAC	240
CTCTCGTGC	ATCCTCGTTC	TAAGACGACC	ACTTATGATT	TCCAGAACTA	CTTGGGGTGT	
TTGGGCATGG	TGATTTTCGC	TTAGTAGCTG	AAC TATTTGA	TTCCACCAGA	ACACATACAA	300
AACCCGTACC	ACTAAAAGCG	AATCATCGAC	TTGATAAACT	AAGGTGGTCT	TGTGTATGTT	
ACAGAAAAGA	GCCAGACATG	AACAAAGTCA	AGCTTTTCTC	AACAGTTGCC	CATGGAAACA	360
TGCTTTTTCT	CGGTCTGTAC	TTGTTTCAGT	TCGAAAAGAG	TTGTCAACGG	GTACCTTTGT	
AAAGTGCAAG	AAGAAAAGCT	TACAATGGTT	CTAGAAGGAA	TATTTTTTCT	CGCCGTTCTT	420
TTTCACGTTT	TTCTTTTCGA	ATGTTACCAA	GATCTTCCTT	ATAAAAAGGA	GCGGCAAGAA	
TTGATAAAAG	AAATACAGAG	GTTACTGAAA	AGCCTGGTGC	CAAGATGTTT	TGGAACAATT	480
AAC TATTTT	TTTATGTCTC	CAATGACTTT	TCGGACCACG	GTTCTACAAG	ACCTTGTTAA	
TTTTGGTTAA	AATGAATGGA	GCCCCACAGA	ATACAAGCCA	TGGCAGTAAA	GCACAGGAAA	540
AAAACCAATT	TTACTTACCT	CGGGGTGTCT	TATGTTCCGT	ACCGTCATTT	CGTGTCTTTT	
TAATGAAAGA	AGCTTGCAAA	ACCTTGTTTT	TCCTCAGAA	TATTGTACAT	GAAAAC TGTG	600
ATTACTTTCT	TCGAACGTTT	TGGAACAAAA	AGTGAGTCTT	ATAACATGTA	CTTTTGACAC	
ACAGGATGGT	GATACAGAAC	AATCTGTGCT	TTGGTAAATG	CATCTCTCTC	CATGTTCCAA	660
TGTCCTACCA	CTATGTCTTG	TTAGACACGA	AACCATTTAC	GTAGAGAGAG	GTACAAGGTT	
ATCAGCAAGA	TCGACGAAAT	ACTTGTTCCC	ATTGCTTGCC	GTCCAAATTT	ACCCTGAACC	720
TAGTCGTTCT	AGCTGCTTTA	TGAACAAGGG	TAACGAACGG	CAGGTTTAAA	TGGGACTTGG	
ACCTGACGCT	GAATTGTACT	GGATCTAAGA	ATGTAGTAAA	GGTTGTCATG	ATGGTAGAGG	780
TGGACTGCGA	CTTAACATGA	CCTAGATTCT	TACATCATTT	CCAACAGTAC	TACCATCTCC	
AATGCACGTG	TGAAGCTCAT	AAGAGCAACT	TCCACCAAAC	TGCACAGTTT	AACATGGATA	840
TTACGTGCAC	ACTTCGAGTA	TTCTCGTTGA	AGGTGGTTTG	ACGTGTCAAA	TTGTACCTAT	
CATCTACTAC	CCTGCACCAT	TAAAGGACTG	CCATACAGTA	TGGAAATGCC	CTTTTGTGTTG	900
GTAGATGATG	GGACGTGGTA	ATTTCTGAC	GGTATGTCAT	ACCTTTACGG	GAAAACAACC	
AATATTTGTT	ACATACTATG	CATCTAAAGC	ATTATGTTGC	CTTCTATTTT	ATATAACCAC	960
TTATAAACAA	TGTATGATAC	GTAGATTTCT	TAATACAACG	GAAGATAAAG	TATATTGGTG	
ATGGAATAAG	GATTGTATGA	ATTATAATTA	ACAAATGGCA	TTTTGTGTAA	CATGCAAGAT	1020
TACCTTATTC	CTAACATACT	TAATATTAAT	TGTTTACCGT	AAAACACATT	GTACGTTCTA	

MSRTRKVDL	LLLAIPGLAL	LLLPNAYCAS	CEPVRIPMCK	SMPWNMTKMP	NHLHHSTQAN	60
AILAIEQFEG	LLTTECSQDL	LFFLCAMYAP	ICTIDFQHEP	IKPCKSV CER	ARAGCEPILI	120
KYRHTWPESL	ACEELPVYDR	GVCISPEAIV	TVEQGTDSMP	DFSMDSNNGN	CGSGREHCKC	180
KPMKATQKTY	LKNYNYVIR	AKVKEVKVVC	HDATAIVEVK	EILKSSLVNI	PKDTVTLYTN	240
SGCLCPQLVA	NEEYIIMGYE	DKERTRLLLV	EGSLAEKWRD	RLAKVKRWD	QKLRRPRKSK	300
DPVAPIPNKN	SNSRQARS					

Figure 3. Deduced amino acid sequence of *Xenopus* frazzled protein. SEQ ID NO:3.

Figure 4. Nucleotide sequence of the full-length frazzled cDNA derived from the *Xenopus* organizer. The sense strand of the DNA on top (5' to 3' direction) and the antisense strand on the bottom line (opposite direction). SEQ ID NO:4.

GAATTCCTT	TCACACAGGA	CTCCTGGCAG	AGGTGAATGG	TTAGCCCTAT	GGATTGGTT	60
CTTAAGGGAA	AGTGTGTCCT	GAGGACCGTC	TCCACTTACC	AATCGGGATA	CCTAAACCAA	
TGTTGATTTT	GACACATGAT	TGATTGCTTT	CAGATAGGAT	TGAAGGACTT	GGATTTTAT	120
ACAATAAAA	CTGTGTACTA	ACTAACGAAA	GTCTATCCTA	ACTTCTGAA	CCTAAAAATA	
CTAATTCTGC	ACTTTTAAAT	TATCTGAGTA	ATTGTTTATT	TTGTATTGGA	TGGGACTAAA	180
GATTAAGACG	TGAAAATTTA	ATAGACTCAT	TAACAAGTAA	AACATAACCT	ACCCTGATTT	
GATAAACTTA	ACTCCTTGCT	TTTGACTTGC	CCATAAACTA	TAAGGTGGGG	TGAGTTGTAG	240
CTATTTGAAT	TGAGGAACGA	AAACTGAACG	GGTATTTGAT	ATTCCACCCC	ACTCAACATC	
TTGCTTTTAC	ATGTGCCAG	ATTTTCCCTG	TATTCCCTGT	ATTCCCTCTA	AAGTAAGCCT	300
AACGAAAATG	TACACGGGTC	TAAAAGGGAC	ATAAGGGACA	TAAGGGAGAT	TTCATTCGGA	
ACACATACAG	GTTGGGCAGA	ATAACAATGT	CTCGAACAAG	GAAAGTGGAC	TCATTACTGC	360
TGTGTATGTC	CAACCCGTCT	TATTGTTACA	GAGCTTGTTT	CTTTCACCTG	AGTAATGACG	
TACTGGCCAT	ACCTGGACTG	GCGCTTCTCT	TATTACCCAA	TGCTTACTGT	GCTTCGTGTG	420
ATGACCGGTA	TGGACCTGAC	CGCGAAGAGA	ATAATGGGTT	ACGAATGACA	CGAAGCACAC	
AGCCTGTGCG	GATCCCCATG	TGCAAATCTA	TGCCATGGAA	CATGACCAAG	ATGCCCAACC	480
TCGGACACGC	CTAGGGGTAC	ACGTTTAGAT	ACGGTACCTT	GTACTGGTTC	TACGGGTTGG	
ATCTCCACCA	CAGCACTCAA	GCCAATGCCA	TCCTGGCAAT	TGAACAGTTT	GAAGGTTTGC	540
TAGAGGTGGT	GTCGTGAGTT	CGGTTACGGT	AGGACCGTTA	ACTTGTCAA	CTTCCAAACG	
TGACCACTGA	ATGTAGCCAG	GACCTTTTGT	TCTTTCTGTG	TGCCATGTAT	GCCCCATTT	600
ACTGGTGAAT	TACATCGGTC	CTGGAAAACA	AGAAAGACAC	ACGGTACATA	CGGGGGTAAA	
GTACCATCGA	TTTCCAGCAT	GAACCAATTA	AGCCTTGCAA	GTCCGTGTGC	GAAAGGGCCA	660
CATGGTAGCT	AAAGGTCGTA	CTTGGTTAAT	TCGGAACGTT	CAGGCACACG	CTTTCOCGGT	
GGGCCGGCTG	TGAGCCCAT	CTCATAAAGT	ACCGGCACAC	TTGGCCAGAG	AGCCTGGCAT	720
CCCGGCCGAC	ACTCGGGTAA	GAGTATTTCA	TGGCCGTGTG	AACCGGTCTC	TCGGACCGTA	
GTGAAGAGCT	GCCCGTATAT	GACAGAGGAG	TCTGCATCTC	CCCAGAGGCT	ATCGTCACAG	780
CACCTTCTCGA	CGGGCATATA	CTGTCTCCTC	AGACGTAGAG	GGGTCTCCGA	TAGCAGTGTC	
TGGAACAAGG	AACAGATTCA	ATGCCAGACT	TCTCCATGGA	TTCAAACAAT	GGAAATTGCG	840
ACCTTGTTCC	TTGTCTAAGT	TACGGTCTGA	AGAGGTACCT	AAGTTTGTTA	CCTTTAACGC	
GAAGCGGCAG	GGAGCACTGT	AAATGCAAGC	CCATGAAGGC	AACCCAAAAG	ACGTATCTCA	900
CTTCGCCGTC	CCTCGTGACA	TTTACGTTTC	GGTACTTCCG	TTGGGTTTTT	TGCATAGAGT	
AGAATAATTA	CAATTATGTA	ATCAGAGCAA	AAGTGAAAGA	GGTGAAAGTG	AAATGCCACG	960
TCTTATTAAT	GTTAATACAT	TAGTCTCGTT	TTCACCTTCT	CCACTTTCAC	TTTACGGTGC	
ACGCAACAGC	AATTGTGGAA	GTAAAGGAGA	TTCTCAAGTC	TTCCCTAGTG	AACATTCCTA	1020
TGCGTTGTGC	TTAACACCTT	CATTTCTCTT	AAGAGTTCAG	AAGGGATCAC	TTGTAAGGAT	

MLLLFRAIPM	LLGLMLVLTQ	DCEIAQYYID	EEPPPGTVIA	VLSQHSIFNT	TDIPATNFRL	60
MKQFNNSLIG	VRES DGQLSI	MERIDREQIC	RQSLHCNLAL	DVVSFSKGHF	KLLNVKVEVR	120
DINDHSPHFP	SEIMHVEVSE	SSSVGTRIPL	EIAIDEDVGS	NSIQNFQISN	NSHFSIDVLT	180
RADGVKYADL	VLMRELDREI	QPTYIMELLA	MDGGVPSLSG	TAVVNIRVLD	FNDNSPVFER	240
STIAVDLVED	APLGYLLEL	HATDDDEGVN	GEIVYGFTSL	ASQEVRLFK	INSRTGSVTL	300
EGQVDFETKQ	TYEFEVQAQD	LGNPLTATC	KTVHILDVN	DNTPAITITP	LTTVNAGVAY	360
IPETATKENF	IALISTTDRA	SGSNGQVRCT	LYGHEHFKLQ	QAYEDSYMIV	TTSTLDRENI	420
AAYSLTVVAE	DLGFPSLTK	KYYTVKVSDE	NDNAPVFSKP	QYEASILENN	APGSYITTVI	480
ARDSDSQNG	KVNYRLVDAK	VMGQSLTTFV	SLDADSGVLR	AVRSLDYEKL	KQLDFEIEAA	540
DNGIPQLSTR	VQLNLRIVDQ	NDNCPVITNP	LLNNGSGEVL	LPISAPQNYL	VFQLKAEDSD	600
EGHNSQLFYT	ILRDPSRLFA	INKESGEVFL	KKQLNSDHSE	DLSIVVAVYD	LGRPSLSTNA	660
TVKFILTDSF	PSNVEVVILQ	PSAEEQHQID	MSIIFI AVL A	GGCALLLLAI	FFVACTCKKK	720
AGEFKQVPEQ	HGTCNEERLL	STPSPQSVSS	SLSQSESCQL	SINTESENC S	VSSNQEQHQQ	780
TGIKHSISVP	SYHTSGWHL D	NCAMSISGHS	HMGHISTKVQ	WAKEIVTSMT	VTILILVENQK	840
RRALSSQCRH	KPVLNTQMNO	QGSDMPITIS	ATESTRVQKM	GTAHCNMKRA	IDCLTL	

Figure 5. Deduced amino acid sequence of the Xenopus PAPC (paraxial protocadherin) protein. It encodes a member of the cadherin family of transmembrane proteins that has dorsalizing activity when constructs are injected into Xenopus embryos. SEQ ID NO:5.

CTGGCAGTGT TACTCTTGAA GGCCAAGTTG ATTTTGAGAC CAAGCAGACT TACGAATTTG	1080
GACCGTCACA ATGAGAACCT CCGGTTCAAC TAAACTCTG GTTCGTCTGA ATGCTTAAAC	
AGGTACAAGC CCAAGATTTG GGCCCCAACC CACTGACTGC TACTTGTAAG GTAAGTGTTC	1140
TCCATGTTTC GGTCTTAAAC CCGGGGTTGG GTGACTGACG ATGAACATTT CATTGACAAG	
ATATACTTGA TGTAATGAT AATACCCAG CCATCACTAT TACCCCTCTG ACTACTGTAA	1200
TATATGAAC ACATTTACTA TTATGGGGTC GGTAGTGATA ATGGGGAGAC TGATGACATT	
ATGCAGGAGT TGCCTATATT CCAGAAACAG CCACAAAGGA GAACTTTATA GCTCTGATCA	1260
TACGTCTCA ACGGATATAA GGTCTTTGTC GGTGTTTCCT CTTGAAATAT CGAGACTAGT	
GCACTACTGA CAGAGCCTCT GGATCTAATG GACAAGTTTC CTGTACTCTT TATGGACATG	1320
CGTGATGACT GTCTCGGAGA CCTAGATTAC CTGTTCAAGC GACATGAGAA ATACCTGTAC	
AGCACTTTAA ACTACAGCAA GCTTATGAGG ACAGTTACAT GATAGTTACC ACCTCTACTT	1380
TCGTGAAAT TGATGTCGTT CGAATACTCC TGTCAATGTA CTATCAATGG TGGAGATGAA	
TAGACAGGGA AAACATAGCA GCGTACTCTT TGACAGTAGT TGCAGAAGAC CTTGGCTTCC	1440
ATCTGTCCCT TTTGTATCGT CGCATGAGAA ACTGTCATCA ACGTCTTCTG GAACCGAAGG	
CCTCATTGAA GACCAAAAAG TACTACACAG TCAAGGTTAG TGATGAGAAT GACAATGCAC	1500
GGAGTAACCT CTGGTTTTTC ATGATGTGTC AGTTCCAATC ACTACTCTTA CTGTTACGTG	
CTGTATTTTC TAAACCCAG TATGAAGCTT CTATTCCTGA AAATAATGCT CCAGGCTCTT	1560
GACATAAAG ATTTGGGGTC ATACTTCGAA GATAAGACCT TTTATTACGA GGTCCGAGAA	
ATATACTAC AGTGATAGCC AGAGACTCTG ATAGTGATCA AAATGGCAA GTAAATTACA	1620
TATATTGATG TCACTATCGG TCTCTGAGAC TATCACTAGT TTTACCGTT CATTTAATGT	
GACTTGTGGA TGCAAAAGTG ATGGGCCAGT CACTAACAAC ATTTGTTTCT CTTGATGCGG	1680
CTGAACACCT ACGTTTTTCAC TACCCGGTCA GTGATTGTTG TAAACAAAGA GAACTACGCC	
ACTCTGGAGT ATTGAGAGCT GTTAGGTCTT TAGACTATGA AAACTTAAA CAACTGGATT	1740
TGAGACCTCA TAACTCTCGA CAATCCAGAA ATCTGATACT TTTTGAATT GTTGACCTAA	
TTGAAATTGA AGCTGCAGAC AATGGGATCC CTCAACTCTC CACTCGCGTT CAACTAAATC	1800
AACCTTAACT TCGACGCTG TTACCCTAGG GAGTTGAGAG GTGAGCGCAA GTTGATTTAG	
TCAGAATAGT TGATCAAAAT GATAATTGCC CTGTGATAAC TAATCCTCTT CTTAATAATG	1860
AGTCTTATCA ACTAGTTTTA CTATTAACGG GACACTATTG ATTAGGAGAA GAATTATTAC	
GCTCGGGTGA AGTTCTGCTT CCCATCAGCG CTCCTCAAAA CTATTTAGTT TTCCAGCTCA	1920
CGAGCCCACT TCAAGACGAA GGGTAGTCGC GAGGAGTTTT GATAAATCAA AAGGTCGAGT	
AAGCCGAGGA TTCAGATGAA GGGCACAAC CCCAGCTGTT CTATACCATA CTGAGAGATC	1980
TTCCGCTCCT AAGTCTACTT CCCGTGTTGA GGGTCGACAA GATATGGTAT GACTCTCTAG	
CAAGCAGATT GTTTGCCATT AACAAAGAAA GTGGTGAAGT GTTCCTGAAA AAACAATTAA	2040
GTTCGTCTAA CAAACGGTAA TTGTTTCTTT CACCACTTCA CAAGGACTTT TTTGTTAATT	
ACTCTGACCA TTCAGAGGAC TTGAGCATAG TAGTTGCAGT GTATGACTTG GGAAGACCTT	2100
TGAGACTGGT AAGTCTCCTG AACTCGTATC ATCAACGTCA CATACTGAAC CCTTCTGGAA	
CATTATCCAC CAATGCTACA GTTAAATTCA TCCTACCGA CTCTTTTCTT TCTAACGTTG	2160
GTAATAGGTG GTTACGATGT CAATTTAAGT AGGAGTGGCT GAGAAAAGGA AGATTGCAAC	

Fig. 6. (Continuation page 2, SEQ ID NO:6).

AAGTCGTTAT	TTTGCAACCA	TCTGCAGAAG	AGCAGCACCA	GATCGATATG	TCCATTATAT	2220
TTACGCAATA	AAACGTTGGT	AGACGTCCTC	TCGTCGTGGT	CTAGCTATAC	AGGTAATATA	
TCATTGCAGT	GCTGGCTGGT	GGTTGTGCTT	TGCTACTTTT	GGCCATCTTT	TTTGTGGCCT	2280
AGTAACGTCA	CGACCGACCA	CCAACACGAA	ACGATGAAAA	CCGGTAGAAA	AAACACCGGA	
GTACTTGTA	AAAGAAAGCT	GGTGAATTTA	AGCAGGTACC	TGAACAACAC	GGAACATGCA	2340
CATGAACATT	TTTCTTTCGA	CCACTTAAAT	TCGTCCATGG	ACTTGTTGTG	CCTTGTACGT	
ATGAAGAACG	CCTGTTAAGC	ACCCCATCTC	CCCAGTCGGT	CTCTTCTTCT	TTGTCTCAGT	2400
TACTTCTTGC	GGACAATTCT	TGGGGTAGAG	GGGTGAGCCA	GAGAAGAAGA	AACAGAGTCA	
CTGAGTCATG	CCAACTCTCC	ATCAATACTG	AATCTGAGAA	TTGCAGCGTG	TCCTCTAACC	2460
GACTCAGTAC	GGTTGAGAGG	TAGTTATGAC	TTAGACTCTT	AACGTCGCAC	AGGAGATTGG	
AAGAGCAGCA	TCAGCAAACA	GGCATAAAGC	ACTCCATCTC	TGTACCATCT	TATCACACAT	2520
TTCTCGTCGT	AGTCGTTTGT	CCGTATTTCT	TGAGGTAGAG	ACATGGTAGA	ATAGTGTGTA	
CTGGTTGGCA	CCTGGACAAT	TGTGCAATGA	GCATAAGTGG	ACATTCTCAC	ATGGGGCACA	2580
GACCAACCGT	GGACCTGTTA	ACACGTTACT	CGTATTCACC	TGTAAGAGTG	TACCCCGTGT	
TTAGTACAAA	GGTACAGTGG	GCAAAGGAGA	TAGTGACTTC	AATGACAGTG	ACTCTGATAC	2640
AATCATGTTT	CCATGTCACC	CGTTTCCTCT	ATCACTGAAG	TTACTGTCAC	TGAGACTATG	
TAGTGGAGAA	TCAGAAAAGA	AGAGCATTGA	GCAGCCAATG	CAGGCACAAG	CCAGTGTCTA	2700
ATCACCTCTT	AGTCTTTTCT	TCTCGTAACT	CGTCGGTTAC	GTCCGTGTTC	GGTCACGAGT	
ATACACAGAT	GAATCAGCAG	GGTTCGACA	TGCCGATAAC	TATTTTCAGCC	ACCGAATCAA	2760
TATGTGTCTA	CTTAGTCGTC	CCAAGGCTGT	ACGGCTATTG	ATAAAGTCGG	TGGCTTAGTT	
CAAGGGTCCA	GAAAATGGGA	ACTGCACATT	GCAATATGAA	AAGGGCTATA	GACTGTCTTA	2820
GTTCCACAGT	CTTTTACCCT	TGACGTGTAA	CGTTATACTT	TTCCCGATAT	CTGACAGAAT	
CTCTGTAGCT	CCTGTATATT	ACAATACCTA	CCATGCAAGA	ATGCCTAACC	TGCACATACC	2880
GAGACATCGA	GGACATATAA	TGTTATGGAT	GGTACGTTCT	TACGGATTGG	ACGTGTATGG	
GAACCATACC	CTTAGAGACC	CTTATTACCA	TATCAATAAT	CCTGTTGCTA	ATCGGATGCA	2940
CTTGGTATGG	GAATCTCTGG	GAATAATGGT	ATAGTTATTA	GGACAACGAT	TAGCCTACGT	
GGCGGAATAT	GAAAGAGATT	TAGTCAACAG	AAGTGCAACG	TTATCTCCGC	AGAGATCGTC	3000
CCGCCTTATA	CTTTCTCTAA	ATCAGTTGTC	TTCACGTTGC	AATAGAGGCG	TCTCTAGCAG	
TAGCAGATAC	CAAGAATTCA	ATTACAGTCC	GCAGATATCA	AGACAGCTTC	ATCCTTCAGA	3060
ATCGTCTATG	GTTCTTAAGT	TAATGTCAGG	CGTCTATAGT	TCTGTCGAAG	TAGGAAGTCT	
AATTGCTACA	ACCTTTTAAT	CATTAGGCAT	GCAAGTGAGA	ATGCACAAAG	GCAAGTGCTT	3120
TTAACGATGT	TGGAAAATTA	GTAATCCGTA	CGTTCACTCT	TACGTGTTTC	CGTTCACGAA	
TAGCATGAAA	GCTAAATATA	TGGAGTCTCC	CCTTTCCCTC	TGATGGATGG	GGGGAGACAC	3180
ATCGTACTTT	CGATTATAT	ACCTCAGAGG	GGAAAGGGAG	ACTACCTACC	CCCCTCTGTG	
AGGACAGTGC	ATAAATATAC	AGCTGCTTTC	TATTTGCATT	TCACTTGGGA	ATTTTTTGTG	3240
TCCTGTCAAG	TATTTATATG	TCGACGAAAG	ATAAACGTAA	AGTGAACCCT	TAAAAACAA	
TTTTTTACAT	ATTTATTTTT	CCTGAATTGA	ATGTGACATT	GTCCTGTGAC	CTAACTAGCA	3300
AAAAAATGTA	TAAATAAAAA	GGACTTAACT	TACACTGTAA	CAGGACAGTG	GATTGATCGT	

Fig. 6. (Continuation page 3, SEQ ID NO:6).

ATTAAATCCA CAGACCTACA GTCAAATATT TGAGGGCCCC TGAAACAGCA CATCAGTCAG	3360
TAATTTAGGT GTCTGGATGT CAGTTTATAA ACTCCCGGGG ACTTTGTCGT GTAGTCAGTC	
GACCTAAAGT GGCCTTTTTA CTTTTCAGAG CTCCTGGGTC TGCCCTCTGT GTTAATCAGC	3420
CTGGATTTC ACGGAAAAAT GAAATCGTC GAGGACCCAG ACGGGAGACA CAATTAGTCG	
CCCTGGTCAA GTCCTGAGTA GGATCATGGC GTTTTATAT GCATCTCACC TACTTTGGAC	3480
GGGACCAGTT CAGGACTCAT CCTAGTACCG CAAAAATATA CGTAGAGTGG ATGAAACCTG	
GTGATTTACA CATAATAGGA AACGCTTGGT TTCAGTGAAG TCTGTGTTGT ATATATTCTG	3540
CACTAAATGT GTATTATCCT TTGCGAACCA AAGTCACTTC AGACACAACA TATATAAGAC	
TTATATACAC GCATTTTGTG TTTGTGTATA TATTTCAGT CCATTCAGAT ATGTGTATAT	3600
AATATATGTG CGTAAACAC AAACACATAT ATAAAGTTCA GGTAAGTCTA TACACATATA	
AGTGCAGACC TTGTAAATTA AATATTCTGA TACTTTTCC TCAATAAATA TTAAAT	
TCACGTCTGG AACATTTAAT TTATAAGACT ATGAAAAAGG AGTTATTTAT AAATTTA	

Fig. 6. (Continuation page 4, SEQ ID NO:6).

FIG. 6. (Continuation page 4, SEQ ID NO:6).

MVCCGPGRML LGWAGLLVLA ALCLLQVPGA QAAACEPVRI PLCKSLPWNM TKMPNHLHHS	60
TQANAILAME QFEGLLGTHC SPDLLFFLCA MYAPICTIDF QHEPIKPCKS VCERARQGCE	120
PILIKYRHSW PESLACDELP VYDRGVCISP EAIVTADGAD FPMDSSTGHC RGASSERCKC	180
KPV RATQKTY FRNNYNYVIR AKVKEVKMKC HDVTAVVEVK EILKASLVNI PRDTVNLVTT	240
SGCLCPPLTV NEEYVIMGYE DEERSRLLLV EGSIAEKWKD RLGKKVKRWD MKLRHLGLGK	300
TDASDSTQNQ KSGRNSNPRP ARS.	

Figure 7. Deduced amino acid sequence of mouse FRZB-1 protein. SEQ ID NO:7.

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Figure 8. Nucleotide sequence of the full-length mouse FRZB-1 cDNA. SEQ ID NO:8.

AAGCCTGGGA CCATGGTCTG CTGCGGCCCG GGACGGATGC TGCTAGGATG GGCCGGGTG	60
TTCGGACCCT GGTACCAGAC GACGCCGGGC CCTGCCTACG ACGATCCTAC CCGGCCCAAC	
CTAGTCCTGG CTGCTCTCTG CCTGCTCCAG GTGCCCGGAG CTCAGGCTGC AGCCTGTGAG	120
GATCAGGACC GACGAGAGAC GGACGAGGTC CACGGGCCTC GAGTCCGACG TCGGACACTC	
CCTGTCCGCA TCCCGCTGTG CAAGTCCCTT CCCTGGAACA TGACCAAGAT GCCCAACCAC	180
GGACAGGCGT AGGGCGACAC GTTCAGGGAA GGGACCTTGT ACTGGTTCTA CGGGTTGGTG	
CTGCACCACA GCACCCAGGC TAACGCCATC CTGGCCATGG AACAGTTCGA AGGGCTGCTG	240
GACGTGGTGT CGTGGGTCCG ATTGCGGTAG GACCGGTACC TTGTCAAGCT TCCCAGACGAC	
GGCACCCACT GCAGCCCGGA TCTTCTCTTC TTCCTCTGTG CAATGTACGC ACCCATTTGC	300
CCGTGGGTGA CGTCGGGCCT AGAAGAGAAG AAGGAGACAC GTTACATGCG TGGGTAAACG	
ACCATCGACT TCCAGCACGA GCCCATCAAG CCCTGCAAGT CTGTGTGTGA GCGCGCCCGA	360
TGGTAGCTGA AGGTCGTGCT CGGGTAGTTC GGGACGTTCA GACACACACT CGCGCGGGCT	
CAGGGCTGCG AGCCCATTTCT CATCAAGTAC CGCCACTCGT GGCCGAAAG CTTGGCCTGC	420
GTCCCGACGC TCGGGTAAGA GTAGTTCATG GCGGTGAGCA CCGGCCTTTC GAACCGGACG	
GACGAGCTGC CGGTGTACGA CCGCGGCGTG TGCATCTCTC CTGAGGCCAT CGTCACCGCG	480
CTGCTCGACG GCCACATGCT GGCGCCGCAC ACGTAGAGAG GACTCCGGTA GCAGTGGCGC	
GACGGAGCGG ATTTTCCTAT GGATTCAAGT ACTGGACACT GCAGAGGGGC AAGCAGCGAA	540
CTGCCTCGCC TAAAAGGATA CCTAAGTTCA TGACCTGTGA CGTCTCCCCG TTCGTCGCTT	
CGTTGCAAAT GTAAGCCTGT CAGAGCTACA CAGAAGACCT ATTTCCGGAA CAATTACAAC	600
GCAACGTTTA CATTCGGACA GTCTCGATGT GTCTTCTGGA TAAAGGCCTT GTTAATGTTG	
TATGTCATCC GGGCTAAAGT TAAAGAGGTA AAGATGAAAT GTCATGATGT GACCGCCGTT	660
ATACAGTAGG CCCGATTTC AATTCTCCAT TTCTACTTTA CAGTACTACA CTGGCGGCAA	
GTGGAAGTGA AGGAAATTCT AAAGGCATCA CTGGTAAACA TTCCAAGGGA CACCGTCAAT	720
CACCTTCACT TCCTTTAAGA TTTCCGTAGT GACCATTTGT AAGGTTCCCT GTGGCAGTTA	
CTTTATACCA CCTCTGGCTG CCTCTGTCCT CCACCTACTG TCAATGAGGA ATATGTCATC	780
GAAATATGGT GGAGACCGAC GGAGACAGGA GGTGAATGAC AGTTACTCCT TATACAGTAG	
ATGGGCTATG AAGACGAGGA ACGTTCCAGG TTA CTCTTGG TAGAAGGCTC TATAGCTGAG	840
TACCCGATAC TTCTGCTCCT TGCAAGGTCC AATGAGAACC ATCTTCCGAG ATATCGACTC	
AAGTGGAAGG ATCGGCTTGG TAAGAAAGTC AAGCGCTGGG ATATGAAACT CCGACACCTT	900
TTCACCTTCC TAGCCGAACC ATTCTTTTCA TTCGCGACCC TATACTTTGA GGCTGTGGAA	
GGACTGGGTA AAAGTATGATC TAGCGATTCC ACTCAGAATC AGAAGTCTGG CAGGAATCTC	960
CCTGACCCAT TTTGACTACG ATCGCTAAGG TGAGTCTTAG TCTTCAGACC GTCCTTGAGA	

AATCCCCGGC	CAGCACGCAG	CTAAATCCTG	AAATGTAAAA	GGCCACACCC	ACGGACTCCC	1020
TTAGGGGCCG	GTCGTGCGTC	GATTTAGGAC	TTTACATTTT	CCGGTGTGGG	TGCCTGAGGG	
TTCTAAGACT	GGCGCTGGTG	GACTAACAAA	GGAAAACCGC	ACAGTTGTGC	TCGTGACCGA	1080
AAGATTCTGA	CCGCGACCAC	CTGATTGTTT	CCTTTTGGCG	TGTCAACACG	AGCACTGGCT	
TTGTTTACCG	CAGACACCGC	GTGGCTACCG	AAGTTACTTC	CGGTCCCCTT	TCTCCTGCTT	1140
AACAAATGGC	GTCTGTGGCG	CACCGATGGC	TTCAATGAAG	GCCAGGGGAA	AGAGGACGAA	
CTTAATGGCG	TGGGGTTAGA	TCCTTTAATA	TGTTATATAT	TCTGTTTCAT	CAATCACGTG	1200
GAATTACCGC	ACCCCAATCT	AGGAAATTAT	ACAATATATA	AGACAAAAGTA	GTTAGTGCAC	
GGGACTGTTC	TTTTGCAACC	AGAATAGTAA	ATTAAATATG	TTGATGCTAA	GGTTTCTGTA	1260
CCCTGACAAG	AAAACGTTGG	TCTTATCATT	TAATTTATAC	AACTACGATT	CCAAAGACAT	
CTGGACTCCC	TGGGTTTAAT	TTGGTGTTCT	GTACCCTGAT	TGAGAATGCA	ATGTTTCATG	1320
GACCTGAGGG	ACCCAAATTA	AACCACAAGA	CATGGGACTA	ACTCTTACGT	TACAAAGTAC	
TAAAGAGAGA	ATCCTGGTCA	TATCTCAAGA	ACTAGATATT	GCTGTAAGAC	AGCCTCTGCT	1380
ATTTCTCTCT	TAGGACCAGT	ATAGAGTTCT	TGATCTATAA	CGACATTCTG	TCGGAGACGA	
GCTGCGCTTA	TAGTCTTG TG	TTTGTATGCC	TTTGTCCATT	TCCCTCATGC	TGTGAAAGTT	1440
CGACGCGAAT	ATCAGAACAC	AAACATACGG	AAACAGGTAA	AGGGAGTACG	ACACTTTCAA	
ATACATGTTT	ATAAAGGTAG	AACGGCATT	TGAAATCAGA	CACTGCACAA	GCAGAGTAGC	1500
TATGTACAAA	TATTTCCATC	TTGCCGTAAA	ACTTTAGTCT	GTGACGTGTT	CGTCTCATCG	
CCAACACCAG	GAAGCATTTA	TGAGGAAACG	CCACACAGCA	TGACTTATTT	TCAAGATTGG	1560
GGTTGTGGTC	CTTCGTAAAT	ACTCCTTTGC	GGTGTGTCGT	ACTGAATAAA	AGTTCTAACC	
CAGGCAGCAA	AATAAATAGT	GTTGGGAGCC	AAGAAAAGAA	TATTTTGCCT	GGTTAAGGGG	1620
GTCCGTCGTT	TTATTTATCA	CAACCCTCGG	TTCTTTTCTT	ATAAAACGGA	CCAATTCCCC	
CACACTGGAA	TCAGTAGCCC	TTGAGCCATT	AACAGCAGTG	TTCTTCTGGC	AAGTTTTTGA	1680
GTGTGACCTT	AGTCATCGGG	AACTCGGTAA	TTGTGTCAC	AAGAAGACCG	TTCAAAAAC	
TTTGTTTCATA	AATGTATTCA	CGAGCATTAG	AGATGAACTT	ATAACTAGAC	ATCTGTTGTT	1740
AAACAAGTAT	TTACATAAGT	GCTCGTAATC	TCTACTTGAA	TATTGATCTG	TAGACAACAA	
ATCTCTATAG	CTCTGCTTCC	TTCTAAATCA	AACCCATTGT	TGGATGCTCC	CTCTCCATTC	1800
TAGAGATATC	GAGACGAAGG	AAGATTTAGT	TTGGGTAACA	ACCTACGAGG	GAGAGGTAAG	

MVCGSPGGML LLRAGLLALA ALCLLRVPGA RAAACEPVRI PLCKSLPWNM TKMPNHLHHS 60
TQANAILAIE QFEGLLGTHC SPDLLFFLCA MYAPICTIDF QHEPIKPCKS VCERARQGCE 120
PILIKYRHSW PENLACEELP VYDRGVCISP EAIVTADGAD FPMDSNGNC RGASSERCKC 180
KPIRATQKTY FRNNYNYVIR AKVKEIKTKC HDVTAVVEVK EILKSSLVNI PRDTVNLYTS 240
SGCLCPPLNV NEEYIIMGYE DEERSRLLLV EGSIAEKWKD RLGKKVKRWD MKLRHLGLSK 300
SDSSNSDSTQ SQKSGRNSNP RQARN.

Figure 9. Deduced amino acid sequence of human FRZB-1 protein. SEQ ID NO:9.

Figure 10. Nucleotide sequence of the full-length human FRZB-1 cDNA. SEQ ID NO:10.

This sequence was assembled from public ESTs from the Genbank database

(accession numbers: H18848, R63748, W38677, W44760, H38379 and N71244).

GGCGGAGCGG GCCTTTTGGC GTCCACTGCG CGGCTGCACC CTGCCCCATC TGCCGGGATC 60
CCGCCTCGCC CGGAAAACCG CAGGTGACGC GCCGACGTGG GACGGGGTAG ACGGCCCTAG
ATGGTCTGCG GCAGCCCCGGG AGGGATGCTG CTGCTGCGGG CCGGGCTGCT TGCCCTGGCT 120
TACCAGACGC CGTCGGGGCCC TCCCTACGAC GACGACGCCC GGCCCCACGA ACGGGACCGA
GCTCTCTGCC TGCTCCGGGT GCCCGGGGCT CGGGCTGCAG CCTGTGAGCC CGTCCGCATC 180
CGAGAGACGG ACGAGGCCCA CGGGCCCCGA GCCCGACGTC GGACACTCGG GCAGGCGTAG
CCCCTGTGCA AGTCCCTGCC CTGGAACATG ACTAAGATGC CCAACCACCT GCACCACAGC 240
GGGGACACGT TCAGGGACGG GACCTTGTA TATTCTACG GGTGGGTGGA CGTGGTGTGC
ACTCAGGCCA ACGCCATCCT GGCCATCGAG CAGTTCGAAG GTCTGCTGGG CACCCACTGC 300
TGAGTCCGGT TCGGTAGGA CCGGTAGCTC GTCAAGCTTC CAGACGACCC GTGGGTGACG
AGCCCCGATC TGCTCTTCTT CCTCTGTGCC ATGTACGCGC CCATCTGCAC CATTGACTTC 360
TCGGGGCTAG ACGAGAAGAA GGAGACACGG TACATGCGCG GGTAGACGTG GTAAGTGAAG
CAGCACGAGC CCATCAAGCC CTGTAAGTCT GTGTGCGAGC GGGCCCGGCA GGGCTGTGAG 420
GTCGTGCTCG GGTAGTTCGG GACATTGAGA CACACGCTCG CCCGGGCCGT CCCGACACTC
CCCATACTCA TCAAGTACCG CCACTCGTGG CCGGAGAACC TGGCCTGCGA GGAGCTGCCA 480
GGGTATGAGT AGTTCATGGC GGTGAGCACC GGCCTCTTGG ACCGGACGCT CCTCGACGGT
GTGTACGACA GGGGCGTGTG CATCTCTCCC GAGGCCATCG TTAAGTGGGA CGGAGCTGAT 540
CACATGCTGT CCGCGCACAC GTAGAGAGGG CTCCGGTAGC AATGACGCCT GCCTCGACTA
TTTCCTATGG ATTCTAGTAA CGGAAACTGT AGAGGGGCAA GCAGTGAACG CTGTAAATGT 600
AAAGGATACC TAAGATCATT GCCTTTGACA TCTCCCCGTT CGTCACTTGC GACATTTACA
AAGCCTATTA GAGCTACACA GAAGACCTAT TTCCGGAACA ATTACAACCTA TGTCATTTCGG 660
TTCGGATAAT CTCGATGTGT CTTCTGGATA AAGGCCCTGT TAATGTTGAT ACAGTAAGCC
GCTAAAGTTA AAGAGATAAA GACTAAGTGC CATGATGTGA CTGCAGTAGT GGAGGTGAAG 720
CGATTTCAAT TTCTCTATTT CTGATTCACG GTACTACACT GACGTCATCA CCTCCACTTC
GAGATTCTAA AGTCCTCTCT GGTAAACATT CCACGGGACA CTGTCAACCT CTATACCAGC 780
CTCTAAGATT TCAGGAGAGA CCATTTGTAA GGTGCCCTGT GACAGTTGGA GATATGGTGC
TCTGGCTGCC TCTGCCCTCC ACTTAATGTT AATGAGGAAT ATATCATCAT GGGCTATGAA 840
AGACCGACGG AGACGGGAGG TGAATTACAA TTACTCCTTA TATAGTAGTA CCCGATACTT

GATGAGGAAC CTACTCCTTG	GTTCAGATT CAAGGTCTAA	ACTCTTGGTG TGAGAACCAC	GAAGGCTCTA CTTCCGAGAT	TAGCTGAGAA ATCGACTCTT	GTGGAAGGAT CACCTTCCTA	900
CGACTCGGTA GCTGAGCCAT	AAAAAGTTAA TTTTTCAATT	GCGCTGGGAT CGCGACCCTA	ATGAAGCTTC TACTTCGAAG	GTCATCTTGG CAGTAGAACC	ACTCAGTAAA TGAGTCATTT	960
AGTGATTCTA TCACTAAGAT	GCAATAGTGA CGTTATCACT	TTCCACTCAG AAGGTGAGTC	AGTCAGAAGT TCAGTCTTCA	CTGGCAGGAA GACCGTCCTT	CTCGAACCCC GAGCTTGGGG	1020
CGGCAAGCAC GCCGTTTCGTG	GCAACTAAAT CGTTGATTTA	CCCGAAATAC GGGCTTTATG	AAAAAGTAAC TTTTTCATTG	ACAGTGGACT TGTCACCTGA	TCCTATTAAG AGGATAATTC	1080
ACTTACTTGC TGAATGAACG	ATTGCTGGAC TAACGACCTG	TAGCAAAGGA ATCGTTTCCT	AAATTGCACT TTTAACGTGA	ATTGCACATC TAACGTGTAG	ATATTCTATT TATAAGATAA	1140
GTTTACTATA CAAATGATAT	AAAATCATGT TTTTAGTACA	GATAACTGAT CTATTGACTA	TATTACTTCT ATAATGAAGA	GTTTCTCTTT CAAAGAGAAA	TGGTTTCTGC ACCAAAGACG	1200
TTCTCTCTTC AAGAGAGAAG	TCTCAACCCC AGAGTTGGGG	TTTGTAATGG AAACATTACC	TTTGGGGGCA AAACCCCCGT	GACTCTTAAG CTGAGAATTC	TATATTGTGA ATATAACACT	1260
GTTTTCTATT CAAAGATAA	TCACTAATCA AGTGATTAGT	TGAGAAAAAC ACTCTTTTTG	TGTTCTTTTG ACAAGAAAAAC	CAATAATAAT GTTATTATTA	AAATTAAACA TTTAATTTGT	1320
TGCTGTTACC ACGACAATGG	AGAGCCTCTT TCTCGGAGAA	TGCTGAGTCT ACGACTCAGA	CCAGATGTTA GGTCTACAAT	ATTTACTTTC TAAATGAAAG	TGCACCCCAA ACGTGGGGTT	1380
TTGGGAATGC AACCCTTACG	AATATTGGAT TTATAACCTA	GAAAAGAGAG CTTTTCTCTC	GTTTCTGGTA CAAAGACCAT	TTACAGAAA AAGTGTCTTT	GCTAGATATG CGATCTATAC	1440
CCTTAAAACA GGAATTTTGT	TACTCTGCCG ATGAGACGGC	ATCTAATTAC TAGATTAAATG	AGCCTTATTT TCGGAATAAA	TTGTATGCCT AACATACGGA	TTTGGGCATT AAACCCGTAA	1500
CTCCTCATGC GAGGAGTACG	TTAGAAAGTT AATCTTTCAA	CCAAATGTTT GGTTTACAAA	ATAAAGGTAA TATTTCCATT	AATGGCAGTT TTACCGTCAA	TGAAGTCAAA ACTTCAGTTT	1560
TGTCACATAG ACAGTGTATC	GCAAAGCAAT CGTTTCGTTA	CAAGCACCAG GTTTCGTGGTC	GAAGTGTTTA CTTCACAAAT	TGAGGAAACA ACTCCTTTGT	ACACCCAAGA TGTGGGTTCT	1620
TGAATTATTT ACTTAATAAA	TTGAGACTGT AACTCTGACA	CAGGAAGTAA GTCCTTCATT	AATAAATAGG TTATTTATCC	AGCTTAAGAA TCGAATTCTT	AGAACATTTT TCTTGTAATAA	1680
GCCTGATTGA CGGACTAACT	GAAGCACAAAC CTTCGTGTTG	TGAAACCACT ACTTTGGTCA	AGCCGCTGGG TCGGCGACCC	GTGTTAATGG CACAATTACC	TAGCATTCTT ATCGTAAGAA	1740
CTTTTGGCAA GAAACCGTT	TACATTTGAT ATGTAAACTA	TTGTTTCATGA AACAAGTACT	ATATATTAAT TATATAATTA	CAGCATTAGA GTCGTAATCT	GAAATGAATT CTTTACTTAA	1800
ATAACTAGAC TATTGATCTG	ATCTGCTGTT TAGACGACAA	ATCACCATAG TAGTGGTATC	TTTTGTTTAA AAAACAAATT	TTTGCTTCCT AAACGAAGGA	TTTAAATAAA AAATTTATTT	1860
CCCATTGGTG GGGTAACCAC	AAAGTCAAAA TTTCAGTTTT	AAAAAAAAAA TTTTTTTTTT	AAA TTT			